SEQUENCE LISTING

<110> Rene S. Hubert

Arthur B. Raitano Daniel E.H. Afar Steve Chappell Mitchell Mary Faris Aya Jakobovits <120> DIAGNOSIS AND THERAPY OF CANCER USING SGP28-RELATED MOLECULES <130> 129.23-US-U1 <150> 60/162,610 <151> 1999-10-28 <160> 36 <170> FastSEQ for Windows Version 4.0 <210> 1 256 <112يم <212> DNA 213> Homo Sapiens ≰400> 1 盤tctctata gtaactgtaa aagtttgaag ctcacattaa cctgtaaaca tcagttggtc 60 agggacagtt gcaaggcctc ctgcaattgt tcaaacagca tttattaaat acgcattaca 120 $c\overline{\underline{\underline{\beta}}}$ ccgagtag ggctatgtag agaggagtca gattatctac ttagatttgg catctactta 180 datttaacat atactagetg agaaattgta ggcatgtttg atacacattt gatttcaaat 240 gëttttcttc tggatc 256 <u>≰2</u>10> 2 211> 2144 &£12> DNA 👯 13> Homo Sapiens ۱. **<4**00> 2 datgaaaca aatacttcat cctgctctgg aaaccactgc aatgacatta ttcccagtgc 60 tgttgttcct ggttgctggg ctgcttccat cttttccaqc aaatqaaqat aaqqatcccq 120 cttttactgc tttgttaacc acccaaacac aagtgcaaag ggagattqtq aataaqcaca atgaactgag gagagcagta teteceeetg ceagaaacat getgaagatg gaatggaaca 240 aagaggctgc agcaaatgcc caaaagtggg caaaccagtg caattacaga cacagtaacc 300 caaaggatcg aatgacaagt ctaaaatgtg gtgagaatct ctacatgtca agtgcccca 360 gctcatggtc acaagcaatc caaagctggt ttgatgagta caatgatttt gactttggtg 420 tagggccaaa gactcccaac gcagtggttg gacattatac acaggttgtt tggtactctt 480 catacctcgt tggatgtgga aatgcctact gtcccaatca aaaaqttcta aaatactact 540 atgtttgcca atattgtcct gctggtaatt gggctaatag actatatgtc ccttatgaac 600 aaggagcacc ttgtgccagt tgcccagata actgtgacga tggactatgc accaatggtt 660 gcaagtacga agatetetat agtaaetgta aaagtttgaa geteacatta aeetgtaaae 720 atcagttggt cagggacagt tgcaaggcat cctgcaattg ttcaaacagc atttattaaa 780 tacgcattac acaccgagta gggctatgta gagaggagtc agattatcta cttagatttg 840 gcatctactt agatttaaca tatactagct gagaaattgt aggcatgttt gatacacatt 900 tgatttcaaa tgtttttctt ctggatctgc tttttatttt acaaaaatat ttttcataca 960 aatggttaaa aagaaacaaa atctataaca acaactttqq atttttatat ataaactttq 1020 tgatttaaat ttactgaatt taattagggt gaaaattttg aaagttgtat tctcatatga 1080 ctaagttcac taaaaccctg gattgaaagt gaaaattatg ttcctagaac aaaatgtaca 1140 aaaagaacaa tataattttc acatgaaccc ttggctgtag ttgcctttcc tagctccact 1200 ctaaggctaa gcatcttcaa agacgttttc ccatatgctg tcttaattct tttcactcat 1260 teaceettet teecaateat etggetggea teeteacaat tgagttgaag etgtteetee 1320



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1740

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2144

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thr Gln Val Gln Arg Glu Ile Val Asn Lys His Asn Glu Leu Arg Arg
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∰a Val Ser Pro Pro Ala Arg Asn Met Leu Lys Met Glu Trp Asn Lys
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Glu Ala Ala Ala Asn Ala Gln Lys Trp Ala Asn Gln Cys Asn Tyr Arg
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trp Phe Asp Glu Tyr Asn Asp Phe Asp Phe Gly Val Gly Pro Lys Thr
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Lys Tyr Tyr Tyr Val Cys Gln Tyr Cys Pro Ala Gly Asn Trp Ala Asn
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Arg Leu Tyr Val Pro Tyr Glu Gln Gly Ala Pro Cys Ala Ser Cys Pro
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